STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/573,600_
Source: _IFwP_
Date Processed by STIC: _4/6/06_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: _10/573,600_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics / Wrapped Aminos — The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2____Invalid Line Length — The rules require that a line **not exceed 72** characters in length. This includes white spaces.

3____Misaligned Amino Numbering — The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.

4____Non-ASCII — The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5____Variable Length — Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0 "bug" — A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7____Skipped Sequences (OLD RULES) — Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped
Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8____Skipped Sequences (NEW RULES) — Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000

9____Use of n's or Xaa's (NEW RULES) — Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10____Invalid <213> Response — Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

11____Use of <220> — Sequence(s) ___ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

12____PatentIn 2.0 "bug" — Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____ Misuse of n/Xaa — "n" can **only** represent a single <u>nucleotide</u>; "Xaa" can **only** represent a single <u>amino acid</u>

AMC – STIC Systems Branch – 03/02/06

IFWP

RAW SEQUENCE LISTING   DATE: 04/06/2006
PATENT APPLICATION: US/10/573,600   TIME: 10:49:35

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Output Set: **N:\CRF4\04062006\J573600.raw**

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 3 <110> APPLICANT: The Trustees of the University of Pennsylvania
 4         Wilson, James M.
 5         Gao, Guangping
 6         Alvira, Mauricio R.
 7         Vandenberghe, Luk H.
 9 <120> TITLE OF INVENTION: Adeno-Associated Virus (AAV) Clades, Sequences, Vectors
10         Containing Same, and Uses Therefor
12 <130> FILE REFERENCE: UPN-P3230PCT
14 <140> CURRENT APPLICATION NUMBER: US/10/573,600
14 <141> CURRENT FILING DATE: 2006-03-24
14 <150> PRIOR APPLICATION NUMBER: US 60/508,226
15 <151> PRIOR FILING DATE: 2003-09-30
17 <160> NUMBER OF SEQ ID NOS: 236
19 <170> SOFTWARE: PatentIn version 3.3
21 <210> SEQ ID NO: 1
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23 <212> TYPE: DNA
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31 aacgctcgag gtcttgtgct tccgggttac aaataccttg gacccggcaa cggactcgac  180
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41 ggaaagaaga ggcctgtaga gcagtctcct caggaaccgg actcctccgc gggtattggc  480
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pp 1-5

Invalid <2137 response

Does Not Comply
Corrected Diskette Needed

see item 10 on Euro summary sheet

RAW SEQUENCE LISTING   DATE: 04/06/2006
PATENT APPLICATION: US/10/573,600   TIME: 10:49:35

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 89 aaaattcctc acacggacgg caactttcac ccttctccgc tgatgggagg gtttggaatg  1920
 91 aagcacccgc ctcctcagat cctcatcaaa aacacacctg tacctgcgga tcctccaacg  1980
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114 aaggggggagc cggtcaacgc agcagacgcg gcggccctcg agcacgacaa ggcctacgac   240
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same error

RAW SEQUENCE LISTING   DATE: 04/06/2006
PATENT APPLICATION: US/10/573,600   TIME: 10:49:35

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186 <213> ORGANISM: adeno-associated virus, human clone 9
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RAW SEQUENCE LISTING   DATE: 04/06/2006
PATENT APPLICATION: US/10/573,600   TIME: 10:49:35

Input Set : A:\UPN-P3230-sequence listing.txt
Output Set: N:\CRF4\04062006\J573600.raw

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367 ggcaagacag gccagcagcc cgcgaaaaag agactcaact ttgggcagac tggcgactca   540
369 gagtcagtgc ccgaccctca accaatcgga gaacccccg caggccctc tggtctggga   600
371 tctggtacaa tggctgcagg cggtggcgct ccaatggcag acaataacga aggcgccgac   660
373 ggagtgggta gttcctcagg aaattggcat tgcgattccg catggctggg cgacagagtc   720
375 atcaccacca gcacccgacc ctgggccctc cccacctaca caaccaccct ctacaagcaa   780
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379 ccctgggggt attttgactt taacagattc cactgccact tctcaccacg tgactggcag   900
381 cgactcatca acaacaactg gggattccgg cccaagagac tcaacttcaa gctcttcaac   960
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428 <212> TYPE: DNA
429 <213> ORGANISM: new AAV serotype, clone hu.41
431 <400> SEQUENCE: 6
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434 gagtggtggg acctgaaacc tggagccccc aagcccaagg ccaaccagca gaagcaggac   120
436 gacggccggg gtctggtgct tcctggctac aagtacctcg gacccttcaa cggactcgac   180
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440 cagcagctca aagcgggtga caatccgtac ctgcggtata accacgccga cgccgagttt   300
442 caggagcgtc tacaagaaga tacgtctttt gggggcaacc tcgggcgagc agtcttccag   360
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```

(handwritten note in right margin:) Please correct this type 1 error in subsequent sequence

VERIFICATION SUMMARY   DATE: 04/06/2006
PATENT APPLICATION: US/10/573,600   TIME: 10:49:36

Input Set : **A:\UPN-P3230-sequence listing.txt**
Output Set: **N:\CRF4\04062006\J573600.raw**

L:14 M:270 C: Current Application Number differs, Replaced Current Application No
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date